| Project Title | Funding | Institution | |
|---|-------------|---|--|
| Simons Simplex Collection support grant | \$10,000 | University of Washington | |
| University of Washington Clinical Site Network Pilot for the National Autism Cohort | \$37,500 | University of Washington | |
| Integrating large scale whole exome data with whole genome data | \$0 | ICAHN SCHOOL OF MEDICINE AT MOUNT SINAI | |
| Sequencing Female-enriched Multiplex Autism Families (FEMFs) | \$0 | Johns Hopkins University | |
| Simons Simplex Collection support grant | \$9,159 | University of Illinois at Chicago | |
| Combining WGS from Utah high-risk pedigrees and SSC families | \$0 | University of Utah | |
| 1/3 - Sequencing Autism Spectrum Disorder Extended Pedigrees | \$298,000 | UNIVERSITY OF UTAH | |
| ASD Family Biobank Program | \$0 | Kaiser Foundation Research Institute | |
| Accelerating Autism Genetics via Whole Population Ascertainment in Denmark | \$0 | Broad Institute, Inc. | |
| Pieces of the Puzzle: Uncovering the Genetics of Autism | \$1,699,790 | Broad Institute, Inc. | |
| Computational tools to analyze SNP data from patients with mental illness | \$572,792 | PARTEK, INC. | |
| Genome Sequencing pilot of Simons Simplex Collection | \$0 | University of Washington | |
| Structural Variation and the Genetic Architecture of Autism | \$0 | University of Washington | |
| Genetic basis of phenotypic variability in 16p11.2 deletion or duplication | \$285,856 | University of Washington | |
| 3/3-Sequencing Autism Spectrum Disorder Extended Pedigrees | \$160,000 | University of Pennsylvania | |
| Autism Spectrum Disorders: Genomes to Outcomes | \$0 | Hospital for Sick Children | |
| Mutations in noncoding DNA and the missing heritability of autism | \$244,030 | University of California, San Diego | |
| Dosage effects of DUF1220 gene subtype CON1 in autism | \$125,000 | University of Colorado, Denver | |
| Extending ASD risk locus discovery to the non-coding genome - Core | \$0 | The Regents of the University of California, San Francisco (Contracts & Grants) | |
| Validation of candidate ASD genes by targeted sequencing with molecular inversion probes | \$101,258 | The Regents of the University of California, San Francisco (Contracts & Grants) | |
| Simons Simplex Collection support grant | \$8,912 | Vanderbilt University | |
| Sequence-based discovery of genes with pleiotropic effects across diagnostic boundaries and throughout the lifespan | \$14,998 | Massachusetts General Hospital | |
| Cryptic Genetic Causes of Autism | \$141,719 | Massachusetts General Hospital | |
| Complex Genetic Architecture of Chromosomal Aberrations in Autism | \$248,999 | Massachusetts General Hospital | |
| Genetic Epidemiology of Complex Traits | \$747,204 | National Institutes of Health | |
| Autism genetics: homozygosity mapping and functional validation | \$765,736 | CHILDREN'S HOSPITAL CORPORATION | |
| Genetic basis of autism | \$4,000,000 | Cold Spring Harbor Laboratory | |
| 2/3 Sequencing Autism Spectrum Disorder Extended Pedigrees | \$231,750 | University of Washington | |
| Whole-exome sequencing to identify causative genes for autism | \$134,203 | ROCKEFELLER UNIVERSITY | |
| Genomic influences on development and outcomes in infants at risk for autism | \$0 | University of Alberta | |
| Genomic profiling of autism families using whole-genome sequencing | \$0 | Institut Pasteur | |

| Project Title | Funding | Institution | |
|--|-------------|---|--|
| Applications of novel statistical methods to CNVs in autism and schizophrenia | \$200,000 | Columbia University | |
| Novel Statistical methods for DNA Sequencing Data, and applications to Autism. | \$318,575 | Columbia University | |
| Simons Simplex Collection support grant | \$13,200 | University of California, Los Angeles | |
| Extending ASD risk locus discovery to the non-coding genome - Project 1 | \$0 | The Trustees of Columbia University in the City of New York | |
| Simons Simplex Collection support grant | \$5,983 | Baylor College of Medicine | |
| Simons Simplex Collection support grant | \$1,831 | Weill Cornell Medical College | |
| Simons Simplex Collection support grant | \$10,000 | University of Missouri | |
| University of North Carolina Clinical Site Network Pilot for the National Autism Cohort | \$24,750 | University of North Carolina | |
| Simons Simplex Collection support grant | \$8,800 | Emory University | |
| Next Generation Gene Discovery in Familial Autism | \$644,823 | University of Washington | |
| Identification and analysis of functional networks perturbed in autism | \$250,000 | Columbia University | |
| Elucidating pathogenic mutations disrupting RNA regulation in autism | \$225,000 | Columbia University | |
| Extending ASD risk locus discovery to the non-coding genome - Project 2 | \$0 | Yale University | |
| Genome-wide analysis of cis-regulatory elements in autism | \$62,500 | Washington University in St. Louis | |
| Simons Simplex Collection support grant | \$10,000 | McGill University Health Centre- Montreal Children's Hospital | |
| New York Genome Center, Inc. | \$2,210,000 | New York Genome Center, Inc. | |
| Illumina, Inc. | \$0 | Illumina, Inc. | |
| MSSNG | \$5,802,895 | Autism Speaks (AS) | |
| Thompson Center Clinical Site Network Pilot for the National Autism Cohort | \$37,500 | The Curators of the University of Missouri | |
| The Future of Genomics Medicine in Patient Care: Contributions from CHOP | \$906,296 | Children's Hospital of Philadelphia | |
| Integrating the genomics of Autism Spectrum Disorders(ASD) in consanguineous and "idiopathic" families | \$665,939 | Yale University | |
| Role of Selfish Spermatogonial Selection in Neurocognitive Disorders | \$59,995 | University of Oxford | |
| FUNCTIONAL DISSECTION OF CNVS IN NEURODEVELOPMENTAL TRAITS | \$366,666 | Duke University | |
| Use of High-throughput Splicing Assays to Prioritize Autism Gene Candidates | \$62,500 | Brown University | |
| Discovery of regulatory variants underlying pediatric neurological disease | \$0 | HudsonAlpha Institute for Biotechnology | |
| High-throughput Screening of Novel Trinucleotide Repeat Expansion in Autism Spectrum Disorders | \$15,000 | The Hospital for Sick Children | |
| DEVELOPING NEW STATISICAL METHODS TO DETECT RARE VARIANTS INVOLVED IN NEUROPSYCHIATRIC DISORDERS | \$433,800 | National Institutes of Health | |
| Simons Simplex Collection support grant | \$10,000 | Yale University | |

| Project Title | Funding | Institution |
|--|-----------|------------------------------------|
| Investigating the role of somatic mutations in autism spectrum disorders | \$263,892 | OREGON HEALTH & SCIENCE UNIVERSITY |
| Identifying Patterns of Genetic Variants Conferring Risk for Neurodevelopmental Disorders | \$29,987 | Pennsylvania State University |